

RAW SEQUENCE LISTING

**The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.**

Application Serial Number: 10/579, 111
Source: IFWP
Date Processed by STIC: 05/23/2006

ENTERED



IFWP

RAW SEQUENCE LISTING

DATE: 05/23/2006

PATENT APPLICATION: US/10/579,111

TIME: 14:26:08

Input Set : A:\21561P SEQLIST.TXT

Output Set: N:\CRF4\05232006\J579111.raw

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4 <110> APPLICANT: Bryan, Janine T.
5     Brownlow, Micelle K.
6     Schultz, Loren D.
7     Wang, Xin-Min
8     Jansen, Kathrin U.
10 <120> TITLE OF INVENTION: OPTIMIZED EXPRESSION OF HPV 58 L1 IN
11     YEAST
13 <130> FILE REFERENCE: 21561P
C--> 15 <140> CURRENT APPLICATION NUMBER: US/10/579,111
C--> 15 <141> CURRENT FILING DATE: 2006-05-11
15 <150> PRIOR APPLICATION NUMBER: PCT/US2004/037372
16 <151> PRIOR FILING DATE: 2004-11-10
18 <150> PRIOR APPLICATION NUMBER: 60/519,211
19 <151> PRIOR FILING DATE: 2003-11-12
21 <160> NUMBER OF SEQ ID NOS: 10
23 <170> SOFTWARE: FastSEQ for Windows Version 4.0
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26 <211> LENGTH: 1497
27 <212> TYPE: DNA
28 <213> ORGANISM: Artificial Sequence
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31 <223> OTHER INFORMATION: HPV58 L1-R
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36 agattgttgg ctgttggttaa cccatacttc tccatcaagt ctccaaacaa caacaagaag 180
37 gtcttggttc caaagggtctc tggtttgcaa tacagagtct tcagagtcag attgccagac 240
38 ccaaacaagt tcggtttccc agacacttcc ttctacaacc cagacactca aagattgggtc 300
39 tgggcttggtg tcggtttgga aatcggtaga ggtcaaccat tgggtgttgg tgtctctggt 360
40 caccatact tcaacaagt tgcgacaccc gaaacctcca acagataccc agctcaacca 420
41 ggttctgaca acagagaatg tttgtccatg gactacaagc aaacccaatt gtgtttgatc 480
42 ggttgtaagc caccaactgg tgaacactgg ggtaaggggtg ttgcttgtaa caacaacgct 540
43 gctgctaccg actgtccacc attggaattg ttcaactcca tcatcgaaga cggtgacatg 600
44 gtcgacactg gtttcggttg tatggacttc ggtaccttgc aagctaacaa gtccgacgtt 660
45 ccaatcgaca tctgtaactc cacctgtaag taccagact acttgaagat ggcttctgaa 720
46 ccatacgggtg actccttggt cttcttcttg agaagagaac aaatgttcgt cagacacttc 780
47 ttcaacagag ctggtaagtt ggggtgaagct gttccagacg acttgtacat caagggttct 840
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49 acctctgaat ctcaattggt caacaagcca tactgggtgc aaagagctca aggtcacaac 960
50 aacggatatct gttggggtaa ccaattgttc gtcactgtcg tcgacaccac tagatccact 1020
51 aacatgacct tgtgtaccga agtcaccaag gaaggtaacct acaagaacga caacttcaag 1080
52 gaatacgtca gacacgtcga ggaatacgac ttgcaattcg tcttccaatt gtgtaagatc 1140
53 accttgactg ctgaaatcat gacctacatc cacaccatgg actctaakat cttggaagac 1200

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55 acctctcaag ctatcacctg tcaaaagact gctccaccaa aggaaaagga agaccattg 1320
56 aacaagtaca ccttctggga agtcaacttg aaggaaaagt tctctgctga cttggaccaa 1380
57 ttcccattgg gtagaaagtt cttgttgcaa tctggtttga aggctaagcc aagattgaag 1440
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61 <211> LENGTH: 498
62 <212> TYPE: PRT
63 <213> ORGANISM: Human Papillomavirus Type 58
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67 1 5 10 15
68 Pro Val Ser Lys Val Val Ser Thr Asp Glu Tyr Val Ser Arg Thr Ser
69 20 25 30
70 Ile Tyr Tyr Tyr Ala Gly Ser Ser Arg Leu Leu Ala Val Gly Asn Pro
71 35 40 45
72 Tyr Phe Ser Ile Lys Ser Pro Asn Asn Asn Lys Lys Val Leu Val Pro
73 50 55 60
74 Lys Val Ser Gly Leu Gln Tyr Arg Val Phe Arg Val Arg Leu Pro Asp
75 65 70 75 80
76 Pro Asn Lys Phe Gly Phe Pro Asp Thr Ser Phe Tyr Asn Pro Asp Thr
77 85 90 95
78 Gln Arg Leu Val Trp Ala Cys Val Gly Leu Glu Ile Gly Arg Gly Gln
79 100 105 110
80 Pro Leu Gly Val Gly Val Ser Gly His Pro Tyr Phe Asn Lys Phe Asp
81 115 120 125
82 Asp Thr Glu Thr Ser Asn Arg Tyr Pro Ala Gln Pro Gly Ser Asp Asn
83 130 135 140
84 Arg Glu Cys Leu Ser Met Asp Tyr Lys Gln Thr Gln Leu Cys Leu Ile
85 145 150 155 160
86 Gly Cys Lys Pro Pro Thr Gly Glu His Trp Gly Lys Gly Val Ala Cys
87 165 170 175
88 Asn Asn Asn Ala Ala Ala Thr Asp Cys Pro Pro Leu Glu Leu Phe Asn
89 180 185 190
90 Ser Ile Ile Glu Asp Gly Asp Met Val Asp Thr Gly Phe Gly Cys Met
91 195 200 205
92 Asp Phe Gly Thr Leu Gln Ala Asn Lys Ser Asp Val Pro Ile Asp Ile
93 210 215 220
94 Cys Asn Ser Thr Cys Lys Tyr Pro Asp Tyr Leu Lys Met Ala Ser Glu
95 225 230 235 240
96 Pro Tyr Gly Asp Ser Leu Phe Phe Phe Leu Arg Arg Glu Gln Met Phe
97 245 250 255
98 Val Arg His Phe Phe Asn Arg Ala Gly Lys Leu Gly Glu Ala Val Pro
99 260 265 270
100 Asp Asp Leu Tyr Ile Lys Gly Ser Gly Asn Thr Ala Val Ile Gln Ser
101 275 280 285
102 Ser Ala Phe Phe Pro Thr Pro Ser Gly Ser Met Val Thr Ser Glu Ser
103 290 295 300
104 Gln Leu Phe Asn Lys Pro Tyr Trp Leu Gln Arg Ala Gln Gly His Asn

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106 Asn Gly Ile Cys Trp Gly Asn Gln Leu Phe Val Thr Val Val Asp Thr
107          325          330          335
108 Thr Arg Ser Thr Asn Met Thr Leu Cys Thr Glu Val Thr Lys Glu Gly
109          340          345          350
110 Thr Tyr Lys Asn Asp Asn Phe Lys Glu Tyr Val Arg His Val Glu Glu
111          355          360          365
112 Tyr Asp Leu Gln Phe Val Phe Gln Leu Cys Lys Ile Thr Leu Thr Ala
113          370          375          380
114 Glu Ile Met Thr Tyr Ile His Thr Met Asp Ser Asn Ile Leu Glu Asp
115 385          390          395          400
116 Trp Gln Phe Gly Leu Thr Pro Pro Pro Ser Ala Ser Leu Gln Asp Thr
117          405          410          415
118 Tyr Arg Phe Val Thr Ser Gln Ala Ile Thr Cys Gln Lys Thr Ala Pro
119          420          425          430
120 Pro Lys Glu Lys Glu Asp Pro Leu Asn Lys Tyr Thr Phe Trp Glu Val
121          435          440          445
122 Asn Leu Lys Glu Lys Phe Ser Ala Asp Leu Asp Gln Phe Pro Leu Gly
123          450          455          460
124 Arg Lys Phe Leu Leu Gln Ser Gly Leu Lys Ala Lys Pro Arg Leu Lys
125 465          470          475          480
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140 agacttttgg ctgttggcaa tccatatttt tccatcaaaa gtcccaataa caataaaaaa 180
141 gtattagtct ccaaggtatc aggcttacag tatagggtct ttaggggtgcg tttacctgat 240
142 cccaataaat ttggttttcc tgatacatct tttataacc ctgatacaca acgtttgggtc 300
143 tgggcatgtg taggccttga aataggtagg ggacagccat tgggtgttgg cgtaagtgggt 360
144 catccttatt tcaataaatt tgatgacact gaaaccagta acagatatcc cgcacagcca 420
145 gggctctgata acaggggaatg cttatctatg gattataaac aaacacaatt atgtttaatt 480
146 ggctgtaaac ctcccactgg tgagcattgg ggtaaagggtg ttgcctgtaa caataatgca 540
147 gctgctactg attgtcctcc attggaactt tttaattcta ttattgagga tgggtgacatg 600
148 gtagatacag ggtttggatg catggacttt ggtacattgc aggctaataa aagtgatgtg 660
149 cctattgata tttgtaacag tacatgcaaa tatccagatt atttaaaaaat ggccagtga 720
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151 tttaataggg ccggaaaact tggcgaggct gtcccggatg acctttatat taaagggtcc 840
152 ggtaatactg cagttatcca aagtagtgca ttttttccaa ctctagtgg ctctatgggt 900
153 acctcagaat cacaattatt taataagcct tattggctac agcgtgcaca aggtcataac 960
154 aatggcattt gctggggcaa tcagttattt gttaccgtag ttgataccac tcgtagcact 1020
155 aatatgacat tatgactga agtaactaag gaaggtagat ataaaaatga taattttaag 1080
156 gaatatgtac gtcattgtga agaataatgac ttacagtttg tttttcagct ttgcaaaatt 1140
157 acactaactg cagagataat gacatatata catactatgg attccaatat tttggaggac 1200

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158 tggcaatttg gtttaacacc tcctccgtct gccagtttac aggacacata tagatttggt 1260
159 acctcccagg ctattacttg ccaaaaaaca gcacccccta aagaaaagga agatccatta 1320
160 aataaatata ctttttgga ggtaaactta aaggaaaagt tttctgcaga tctagatcag 1380
161 tttccttttg gacgaaagt tttattacaa tcaggcctta aagcaaagcc cagactaaaa 1440
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167 <213> ORGANISM: Artificial Sequence
169 <220> FEATURE:
170 <223> OTHER INFORMATION: PCR Primer
172 <400> SEQUENCE: 4
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177 <212> TYPE: DNA
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181 <223> OTHER INFORMATION: PCR Primer
183 <400> SEQUENCE: 5
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188 <212> TYPE: DNA
189 <213> ORGANISM: Artificial Sequence
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192 <223> OTHER INFORMATION: PCR Primer
194 <400> SEQUENCE: 6
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198 <211> LENGTH: 28
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200 <213> ORGANISM: Artificial Sequence
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203 <223> OTHER INFORMATION: PCR Primer
205 <400> SEQUENCE: 7
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209 <211> LENGTH: 34
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211 <213> ORGANISM: Artificial Sequence
213 <220> FEATURE:
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216 <400> SEQUENCE: 8
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221 <212> TYPE: DNA
222 <213> ORGANISM: Artificial Sequence

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 232 <212> TYPE: DNA
 233 <213> ORGANISM: Artificial Sequence
 235 <220> FEATURE:
 236 <223> OTHER INFORMATION: HPV 58 L1 antisense
 238 <400> SEQUENCE: 10
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 240 cagcagaggt gactgcttat gcagagatct tggagataga tgatgatgcg accaaggaga 120
 241 tctaacaacc gacaaccatt gggatgaag aggtagtcca gaggtttgtt gttgttcttc 180
 242 cagaaccaag gtttccagag accaaacgtt atgtctcaga agtctcagtc taacgggtctg 240
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 245 gtgggtatga agttgttcaa gctgctgtgg ctttggaggt tgtctatggg tcgagttggg 420
 246 ccaagactgt tgtctcttac aaacaggtac ctgatgttcg tttgggttaa cacaaactag 480
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VERIFICATION SUMMARY

PATENT APPLICATION: US/10/579,111

DATE: 05/23/2006

TIME: 14:26:09

Input Set : A:\21561P SEQLIST.TXT

Output Set: N:\CRF4\05232006\J579111.raw

L:15 M:270 C: Current Application Number differs, Replaced Current Application No

L:15 M:271 C: Current Filing Date differs, Replaced Current Filing Date